```
Db    144 --ALNDSMINETARDAARVQVASTLSVLVGLFQVGLGLIHFGFVVTYLSEPLVRGYTTAA 201

Qy    212 GLQILISVLKYIFGLTIPSYTGPGSIVFTFIDICKNLPHTNIASLIFALISGAFLVLVKE 271
          :|: :| |||:||| : |:::|| |:::| :::|  || : : ::: | ::|  ||:||
Db    202 AVQVFVSQLKYVFGLHLSSHSGPLSLIYTVLEVCWKLPQSKVGTVVTAAVAGVVLVVVKL 261

Qy    272 LNARYMHKIRFPIPTEMIVVVVATAISGGCKMPKKYHMQIVGEIQRGFPTPVSPVVSQWK 331
          || :   ::  ||| |:: :: || || |  : :: : :|| |  |   ||:|    :
Db    262 LNDKLQQQLPMPIPGELLTLIGATGISYGMGLKHRFEVDVVGNIPAGLVPPVAPNTQLFS 321

Qy    332 DMIGTAFSLAIVSYVINLAMGRTLANKHGYDVDSNQEMIALGCSNFFGSFFKIHVICCAL 391
          ::|:||::|:| : | :::|:  | :||| ||||||::||| ||  |  |:   : |::
Db    322 KLVGSAFTIAVVGFAIAISLGKIFALRHGYRVDSNQELVALGLSNLIGGIFQCFPVSCSM 381

Qy    392 SVTLAVDGAGGKSQVASLCVSLVVMITMLVLGIYLYPLPKSVLGALIAVNLKNSLKQLTD 451
          | :|  :  || ||||    || ::: :: ||   : |||:|| |:| |||| |:||:|
Db    382 SRSLVQESTGGNSQVAGAISSLFILLIIVKLGELFHDLPKAVLAAIIIVNLKGMLRQLSD 441

Qy    452 PYYLWRKSKLDCCIWVVSFLSSFFLSLPYGVAVGVAFSVLVVVFQTQFRNGYALAQVMDT 511
            ||: :: |  ||:|:| ::  |:|  |: | | ||:|:|| :||  :   | || ||
Db    442 MRSLWKANRADLLIWLVTFTATILLNLDLGLVVAVIFSLLLVVVRTQMPHYSVLGQVPDT 501

Qy    512 DIYVNPKTYNRAQDIQGIKIITYCSPLYFANSEIF-----------------RQKVIAK 553
          ||| :   |: |::::|:|:    : :||||:| :                 ::|:: |
Db    502 DIYRDVAEYSEAKEVRGVKVFRSSATVYFANAEFYSDALKQRCGVDVDFLISQKKKLLKK 561

Qy    554 --TVSLQELQQD----FENAPPTDP-------------NNNQTPANGTSVSYITFSPDS 593
            : |::||::     : | |                 :||        ||    | |
Db    562 QEQLKLKQLQKEEKLRKQAASPKGASVSINVNTSLEDMRSNNVEDCKMMQVS----SGDK 617

Qy    594 SSPAQSEPPASAEAPGEPSDMLASVPPFVTFHTLILDMSGVSFVDLMGIKALAKLSSTYG 653
          | :     ::|| : | : |   |  ||:|||:  :|||| : :|:|  :   :
Db    618 MEDATANGQEDSKAP-DGSTLKALGLPQPDFHSLILDLGALSFVDTVCLKSLKNIFHDFR 676

Qy    654 KIGVKVFLVNIHAQVYNDISHGGVFEDGSLECKHVFPSIHDAVLFAQANARDV 706
          :| |:|::   |: | : :  |  | | |:  ||:| |:|||| ||  : | |
Db    677 EIEVEVYMAACHSPVVSQLEAGHFF-DASITKKHLFASVHDAVTFALQHPRPV 728


RESULT   2
Q9BXS9
ID   Q9BXS9      PRELIMINARY;      PRT;   759 AA.
AC   Q9BXS9;
DT   01-JUN-2001 (TrEMBLrel. 17, Created)
DT   01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT   01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
DE   SOLUTE CARRIER FAMILY 26 MEMBER 6.
GN   SLC26A6.
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=21145584; PubMed=11247665;
RA   Waldegger S., Moschen I., Ramirez A., Smith R.J., Ayadi H., Lang F.,
RA   Kubisch C.;
RT   "Cloning and Characterization of SLC26A6, a Novel Member of the Solute
RT   Carrier 26 Gene Family.";
RL   Genomics 72:43-50(2001).
DR   EMBL; AF288410; AAK19153.1; -.
SQ   SEQUENCE   759 AA;  82966 MW;  63CB0B756C9675C6 CRC64;


  Query Match             32.5%;  Score 1258;  DB 4;  Length 759;
  Best Local Similarity   36.6%;  Pred. No. 1e-70;
  Matches  261;  Conservative  162;  Mismatches  238;  Indels   52;  Gaps    8;

Qy      32 PVGEKLRNAFRCSSAKIKAVVFGLLPVLSWLPKYKIKDYIIPDLLGGLSGGSIQVPQGMA 91
           |   : |   :|| |:  |::   |||| |||:| ::|::: ||| |||   :|:||||
Db      51 PRTHQWRTWLQCSRARAYALLLQHLPVLVWLPRYPVRDWLLGDLLSGLSVAIMQLPQGLA 110

Qy      92 FALLANLPAVNGLYSSFFPLLTYFFLGGVHQMVPGTFAVISILVGNICLQLAPESKFQVF 151
```

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          :|||| || | ||||||:|:  ||  |    : |||||:|::||::   |||:
Db    111 YALLAGLPPVFGLYSSFYPVFIYFLFGTSRHISVGTFAVMSVMVGSVTESLAPQ------ 164

Qy    152 NNATNESYVDTAAMEAERLHVSATLACLTAIIQMGLGFMQFGFVAIYLSESFIRGFMTAA 211
            | |:| ::  | :| |: |::||: |  : |:||| : |||| |||| :||: |||
Db    165 --ALNDSMINETARDAARVQVASTLSVLVGLFQVGLGLIHFGFVVTYLSEPLVRGYTTAA 222

Qy    212 GLQILISVLKYIFGLTIPSYTGPGSIVFTFIDICKNLPHTNIASLIFALISGAFLVLVKE 271
          :|: :| |||:||| : |:::|| |:::| :::|  || : : ::: | ::|  ||:||
Db    223 AVQVFVSQLKYVFGLHLSSHSGPLSLIYTVLEVCWKLPQSKVGTVVTAAVAGVVLVVVKL 282

Qy    272 LNARYMHKIRFPIPTEMIVVVVATAISGGCKMPKKYHMQIVGEIQRGFPTPVSPVVSQWK 331
          || :   ::  ||| |:: :: || || |  : :: : :|| |  |   ||:|    :
Db    283 LNDKLQQQLPMPIPGELLTLIGATGISYGMGLKHRFEVDVVGNIPAGLVPPVAPNTQLFS 342

Qy    332 DMIGTAFSLAIVSYVINLAMGRTLANKHGYDVDSNQEMIALGCSNFFGSFFKIHVICCAL 391
          ::|:||::|:| : | :::|:  | :||| ||||||::||| ||  |  |:   : |::
Db    343 KLVGSAFTIAVVGFAIAISLGKIFALRHGYRVDSNQELVALGLSNLIGGIFQCFPVSCSM 402

Qy    392 SVTLAVDGAGGKSQVASLCVSLVVMITMLVLGIYLYPLPKSVLGALIAVNLKNSLKQLTD 451
          | :|  :  || ||||    || ::: :: ||   : |||:|| |:| |||| |:||:|
Db    403 SRSLVQESTGGNSQVAGAISSLFILLIIVKLGELFHDLPKAVLAAIIIVNLKGMLRQLSD 462

Qy    452 PYYLWRKSKLDCCIWVVSFLSSFFLSLPYGVAVGVAFSVLVVVFQTQFRNGYALAQVMDT 511
            ||: :: |  ||:|:| ::  |:|  |: | | ||:|:|| :||  :   | || ||
Db    463 MRSLWKANRADLLIWLVTFTATILLNLDLGLVVAVIFSLLLVVVRTQMPHYSVLGQVPDT 522

Qy    512 DIYVNPKTYNRAQDIQGIKIITYCSPLYFANSEIF-----------------RQKVIAK 553
          ||| :   |: |::::|:|:    : :||||:| :                 ::|:: |
Db    523 DIYRDVAEYSEAKEVRGVKVFRSSATVYFANAEFYSDALKQRCGVDVDFLISQKKKLLKK 582

Qy    554 --TVSLQELQQD----FENAPPTDP-------------NNNQTPANGTSVSYITFSPDS 593
            : |::||::     : | |                 :||        ||    | |
Db    583 QEQLKLKQLQKEEKLRKQAASPKGASVSINVNTSLEDMRSNNVEDCKMMQVS----SGDK 638

Qy    594 SSPAQSEPPASAEAPGEPSDMLASVPPFVTFHTLILDMSGVSFVDLMGIKALAKLSSTYG 653
          | :     ::|| : | : |   |  ||:|||:  :|||| : :|:|  :   :
Db    639 MEDATANGQEDSKAP-DGSTLKALGLPQPDFHSLILDLGALSFVDTVCLKSLKNIFHDFR 697

Qy    654 KIGVKVFLVNIHAQVYNDISHGGVFEDGSLECKHVFPSIHDAVLFAQANARDV 706
          :| |:|::   |: | : :  |  | | |:  ||:| |:|||| ||  : | |
Db    698 EIEVEVYMAACHSPVVSQLEAGHFF-DASITKKHLFASVHDAVTFALQHPRPV 749


RESULT   3
Q9EPH0
ID   Q9EPH0      PRELIMINARY;      PRT;   744 AA.
AC   Q9EPH0;
DT   01-MAR-2001 (TrEMBLrel. 16, Created)
DT   01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
DT   01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
DE   PRESTIN.
OS   Rattus norvegicus (Rat).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX   NCBI_TaxID=10116;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Ludwig J., Oliver D., Frank G., Kloecker N., Gummer A.W., Fakler B.;
RT   "The reciprocal electromechanical properties of rat Prestin: the motor
RT   molecule of rat outer hair cells.";
RL   Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AJ303372; CAC21555.1; -.
DR   InterPro; IPR002645; STAS.
DR   InterPro; IPR001902; Sulfate_transp.
DR   Pfam; PF01740; STAS; 1.
DR   Pfam; PF00916; Sulfate_transp; 1.
SQ   SEQUENCE   744 AA;  81278 MW;  E49E842CF7A3CD58 CRC64;


  Query Match             32.5%;  Score 1255.5;  DB 11;  Length 744;
  Best Local Similarity   35.4%;  Pred. No. 1.4e-70;
  Matches  265;  Conservative  163;  Mismatches  267;  Indels   53;  Gaps   11;
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Qy     542 NSEIFRQKVIAKTVSLQELQQDFENAPPTDPNNNQTPANGTSVSYITFSPDSSSPAQSEP 601
           | | |:  :  :||:   ::  :: |             |          | |   |
Db     358 nkecfksalykqtvnpilikvawkkaakrkikekvvtlggiq--------demsvqlshd 409

Qy     602 PASAEAPGEPSDMLASVPPFVTFHTLILDMSGVSFVDLMGIKALAKLSSTYGKIGVKVFL 661
           |                  :  ||:::| | : |:|  ||  | ::   |  ||::| |
Db     410 p------------------lelhtividcsaiqfidtagihtlkevrrdyeaigiqvll 450

Qy     662 VNIHAQVYNDISHGGVFEDGSLECKH-----VFPSIHDAVLFAQAN 702
               :  | : :::|         ||      :| |:::|: ||: :
Db     451 aqcnptvrdsltngey-------ckkeeenllfysvyeamafaevs 489


RESULT   8
AAM42394
ID   AAM42394 standard; Protein; 143 AA.
XX
AC   AAM42394;
XX
DT   22-OCT-2001  (first entry)
XX
DE   Human polypeptide SEQ ID NO 127.
XX
KW   Human; nootropic; neuroprotective; cytostatic; dermatological; virucide;
KW   immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary;
KW   antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer;
KW   antirheumatic; hepatotropic; cerebroprotective; antiinflammatory;
KW   antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal;
KW   antiparasitic; cardiant;  immune disorder; cardiovascular disorder;
KW   neurological disease; infection; nephrotropic; gene therapy; vaccine.
XX
OS   Homo sapiens.
XX
PN   WO200155449-A1.
XX
PD   02-AUG-2001.
XX
PF   17-JAN-2001; 2001WO-US01346.
XX
PR   31-JAN-2000; 2000US-0179065.
PR   04-FEB-2000; 2000US-0180628.
PR   19-MAY-2000; 2000US-0205515.
PR   07-JUL-2000; 2000US-0216880.
PR   14-JUL-2000; 2000US-0218290.
PR   14-AUG-2000; 2000US-0225447.
PR   01-SEP-2000; 2000US-0229343.
PR   06-SEP-2000; 2000US-0230437.
PR   08-SEP-2000; 2000US-0231243.
PR   25-SEP-2000; 2000US-0234997.
PR   29-SEP-2000; 2000US-0236367.
PR   13-OCT-2000; 2000US-0239937.
PR   08-NOV-2000; 2000US-0246476.
PR   08-NOV-2000; 2000US-0246477.
PR   08-NOV-2000; 2000US-0246525.
PR   08-NOV-2000; 2000US-0246526.
PR   08-NOV-2000; 2000US-0246528.
PR   17-NOV-2000; 2000US-0249210.
PR   17-NOV-2000; 2000US-0249211.
PR   17-NOV-2000; 2000US-0249214.
PR   17-NOV-2000; 2000US-0249265.
PR   01-DEC-2000; 2000US-0250160.
PR   01-DEC-2000; 2000US-0250391.
PR   05-DEC-2000; 2000US-0251030.
PR   05-DEC-2000; 2000US-0251988.
PR   05-DEC-2000; 2000US-0256719.
PR   06-DEC-2000; 2000US-0251479.
PR   08-DEC-2000; 2000US-0251989.
PR   08-DEC-2000; 2000US-0251990.
PR   11-DEC-2000; 2000US-0254097.
XX
PA   (HUMA-) HUMAN GENOME SCI INC.
XX
PI   Rosen CA,  Barash SC,  Ruben SM;
XX
DR   WPI; 2001-476225/51.
DR   N-PSDB; AAI62799.
XX
PT   Novel plasma membrane associated proteins useful for diagnosing,
PT   treating, preventing and/or prognosing disorders related to the
PT   proteins, including cancer, immune response and neuronal disorders  -
XX
PS   Claim 11; SEQ ID NO 127; 532pp + Sequence Listing; English.
XX
CC   The invention relates to novel genes (AAI62752-AAI62961) and proteins
CC   (AAM42347-AAM42415) useful for preventing, treating or ameliorating
CC   medical conditions e.g. by protein or gene therapy. The genes are
CC   isolated from a range of human tissues disclosed in the specification.
CC   The nucleic acids, proteins, antibodies and (ant)agonists are useful
CC   in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast
CC   and ovarian cancer and other cancers of the adrenal gland, bone, bone
CC   marrow, breast, gastrointestinal tract, liver, lung, or urogenital;
CC   (b) immune disorders e.g. Addison's disease, allergies, autoimmune
CC   haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
CC   disease, multiple sclerosis, rheumatoid arthritis and ulcerative
CC   colitis; (c) cardiovascular disorders such as myocardial ischaemias;
CC   (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and
CC   epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
CC   and parasitic infections.
CC   Note: The sequence data for this patent did not form part of the
CC   printed specification, but was obtained in electronic format directly
CC   from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ   Sequence   143 AA;


  Query Match             15.5%;  Score 601.5;  DB 22;  Length 143;
  Best Local Similarity   91.0%;  Pred. No. 8.9e-53;
  Matches  122;  Conservative    3;  Mismatches    8;  Indels    1;  Gaps    1;

Qy     107 SFFPLLTYFFLGGVHQMVPGTFAVISILVGNICLQLAPESKFQVFNNATNESYVDTAAME 166
           |||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db       9 sffplltyffxggvhqmvpgtfavisilvgniclqlapeskfqvfnnatnesyvdtaame 68

Qy     167 AERLHVSATLACLTAIIQMGLGFMQFGFVAIYLSESFIRGFMTAAGLQILISVLKYIFGL 226
           |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db      69 aerlhvsatlacltxiiqmglgfmqfgfvaiylsesfirgfmtaaglqilisvlkyifga 128

Qy     227 -TIPSYTGPGSIVF 239
            |::: | || |
Db     129 gqepthswprsiaf 142


RESULT   9
AAY44945
ID   AAY44945 standard; Protein; 593 AA.
XX
AC   AAY44945;
XX
DT   23-MAY-2000  (first entry)
XX
DE   Wheat sulphate permease-2.
XX
KW   Sulphate Permease; sulphate assimilation protein; wheat; probe;
KW   mapping; marker; plant breeding; chimeric gene; transgenic plant;
KW   antibody; screen.
XX
OS   Triticum aestivum.
XX
PN   WO200004154-A2.
XX
PD   27-JAN-2000.
XX
PF   13-JUL-1999;   99WO-US15810.
XX
PR   14-JUL-1998;   98US-0092833.
XX
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